SEQUENCE LISTING

<110> National Agricultural Research Organization; Katsuji Watanabe

<120> METHOD AND SYSTEM FOR SEARCHING FOR RELATIONSHIPS BETWEEN BASE SEQUENCE IN GENES

<160>4

<210>1

<211>1009

<212>DNA

<213>Rhodospirillum salexigens

<220>

<221>rRNA

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<223>unknown

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121 ggaaacggca gctaataccg gatacgccgt attgggaaag aaattcggcc ttggatgggc

181 ccgcgttgga ttagctagat ggtggggtaa cggcctacca tggcgacgat ccatagctgg

241 tttgagagga tgatcagcca cactgggact gagacacggc ccagactcct acgggaggca

301 gcagtgggga atcttagaca atgggggcaa ccctgatcta gccatgccgc gtgagtgatg

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721 ggtagtccac gccgtaaacg atgggtgcta gatgtcgggg ctcttagagt ttcggtatcg

781 cagctaacgc attaagcacc ccgccngggg agtacggccg caaggttaaa actcaaagga

841 attgacgggg gcnngcacaa gcggtggagc atgtggttta attcgaanna acgcgcagaa

901 ccttaccagc tcttgacatc ccgggacgac ttccagagat ggatttttc acttcggtga

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121 tcagggctct gggataactg ctggaaacgg cagctaatac cggatacgcc gtattgggaa

181 agaaattcgg ccttggatgg gcccgcgttg gattagctag atggtggggt aacggcctac

241 catggcgacg atccatagct ggtttgagag gatgatcagc cacactggga ctgagacacg

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481 ggagngggcn agcgttgttc ggaattactg ggcgtaaagc gcgcgtaggc ggatcggtca

541 gttgggggtg aaagcccggg gctcaacctc ggaactgccc tcaaaactac cgatcnagag

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1021 tcgtgagatg ttgggttaag tcccncaacg agcgcaaccc tcgcccttag ttgccagcat

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1261 caactcgaga gcatgaaggt ggaatcgcta gtaatcgcgg atcagcatgc cgcggtgaat

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